



OIPE

Unlabeled
UnlabeledRAW SEQUENCE LISTING
PATENT APPLICATION: US/10/056,744BDATE: 08/22/2002
TIME: 16:34:46Input Set : A:\CORRECTEDSEQUENCE LISTING.txt
Output Set: N:\CRF4\08222002\J056744B.raw

4 <110> APPLICANT: Kapeller-Libermann, Rosana
6 <120> TITLE OF INVENTION: 58860, A HUMAN CHOLESTERYL ESTER
7 HYDROLASE AND USES THEREFOR
9 <130> FILE REFERENCE: MPI2001-026P1RNM
11 <140> CURRENT APPLICATION NUMBER: 10/056,744B
C--> 13 <141> CURRENT FILING DATE: 2002-08-16
15 <150> PRIOR APPLICATION NUMBER: 60/264,167
17 <151> PRIOR FILING DATE: 2001-01-25
19 <160> NUMBER OF SEQ ID NOS: 6
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

366 <210> SEQ ID NO: 6
367 <211> LENGTH: 10
368 <212> TYPE: PRT
369 <213> ORGANISM: Homo sapiens
371 <220> FEATURE:
372 <221> NAME/KEY: VARIANT
373 <222> LOCATION: (1)...(1)
374 <223> OTHER INFORMATION: Xaa = L, I or V
376 <223> OTHER INFORMATION: The amino acid at position 2 can be any amino acid
W--> 378 <221> NAME/KEY: VARIANT
W--> 379 <222> LOCATION: (3)...(3)
W--> 380 <223> Xaa = L, I, V, M, S or T
W--> 382 <221> NAME/KEY: VARIANT
W--> 383 <222> LOCATION: (4)...(4)
W--> 384 <223> Xaa = L, I, V, M, S or T
W--> 386 <221> NAME/KEY: VARIANT
W--> 387 <222> LOCATION: (6)...(6)
W--> 388 <223> Xaa = H, Y, W or V
390 <223> OTHER INFORMATION: The amino acid at position 8
391 can be any amino acid
W--> 393 <221> NAME/KEY: VARIANT
W--> 394 <222> LOCATION: (10)...(10)
W--> 395 <223> Xaa = G, T, A or C
397 <400> SEQUENCE: 6
W--> 398 Xaa Xaa Xaa Xaa Gly Xaa Ser Xaa Gly Xaa
399 1 5 10
E--> 408 - 1 -

8/22/02

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/056,744B

DATE: 08/22/2002

TIME: 16:34:47

Input Set : A:\CORRECTEDSEQUENCE LISTING.txt

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:378 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:379 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:380 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:382 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:383 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:384 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:386 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:387 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:388 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:393 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:394 M:281 W: Numeric Fields not Ordered, <222> Sort in ascending order!
L:395 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:408 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6